Data Visualization in R

Kevin Johnson

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1 Introduction

Ggplot2 is widely considered the best R package for visualizing data. It has a unique way of implementing plots so it can have a steep learning curve, but once you get it the syntax is very powerful.

The core function used to start every plot is ggplot(). The next most important function is aes() which stands for aesthetics. Plots are built by adding layers onto the base function that specify what you want the plot to show (you literally use the + operator to add layers). Parameters passed to the aes() function are passed on to every layer in the plot.

2 Scatterplots

We'll start with something easy, a simple scatterplot. Ggplot2 comes with several datasets preloaded, one of which is called mtcars. It has data on cars extracted from the 1974 Motor Trend US magazine. The variables included are:

- mpg: Miles/(US) gallon
- cyl: Number of cylinders
- disp: Displacement (cu.in.)

• hp: Gross horsepower

• drat: Rear axle ratio

• wt: Weight (lb/1000)

• qsec: 1/4 mile time

• vs: V/S

• am: Transmission (0 = automatic, 1 = manual)

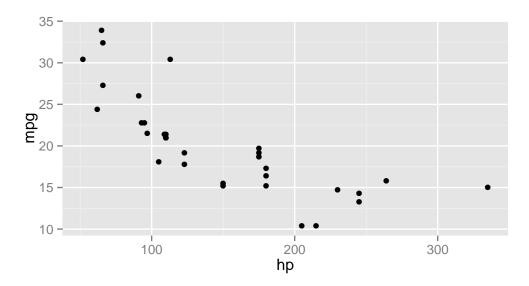
• gear: Number of forward gears

• carb: Number of carburetors

```
head(mtcars)
##
                    mpg cyl disp hp drat
                                            wt qsec vs am gear carb
## Mazda RX4
                   21.0
                          6 160 110 3.90 2.620 16.46 0
                                                                   4
## Mazda RX4 Wag
                          6 160 110 3.90 2.875 17.02
                   21.0
                                                                   4
## Datsun 710
                    22.8
                          4 108 93 3.85 2.320 18.61 1
                                                                   1
## Hornet 4 Drive
                   21.4
                          6 258 110 3.08 3.215 19.44 1
                                                              3
                                                                   1
## Hornet Sportabout 18.7
                          8 360 175 3.15 3.440 17.02 0
                                                              3
                                                                   2
## Valiant
                    18.1
                          6 225 105 2.76 3.460 20.22 1 0
                                                              3
                                                                   1
```

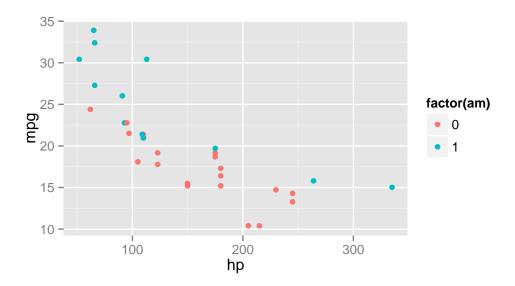
The geom_point() function takes in x and y values and creates a scatterplot of the points. Let's plot the horsepower of the car on the x-axis and the miles per gallon on the y-axis.

```
ggplot(data = mtcars, aes(x = hp, y = mpg)) +
    geom_point()
```

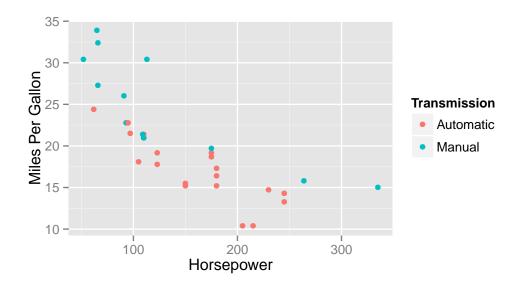


The geom_point() function can take several parameters that determine what the points will look like. You can control things like color, transparency, size, and shape. One of the best features of ggplot2 is that you can set easily set these aesthetic properties to be based on variables in your dataset. For example, let's say we want to have the points be different colors based on if the car is manual or automatic (stored in am).

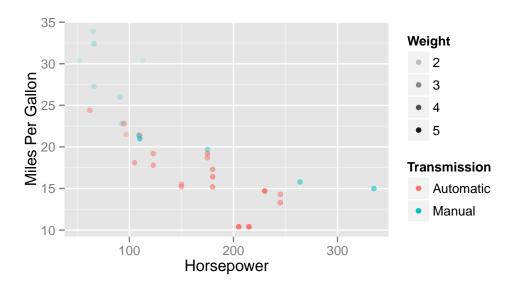
```
ggplot(data = mtcars, aes(x = hp, y = mpg, color = factor(am))) +
    geom_point()
```



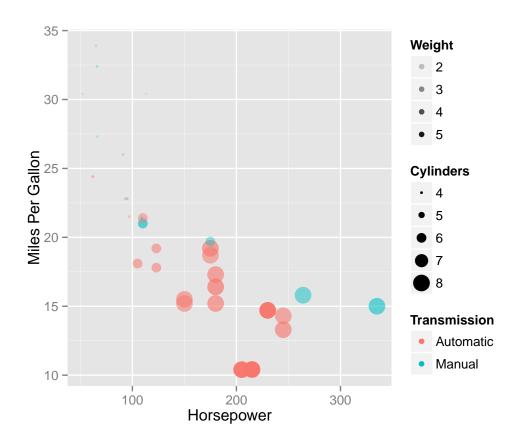
I'll talk about themeing graphs in more detail later, but for now let's just look at how to handle legends and scales in ggplot2. Obviously we don't want our legend to be titled "factor(am)", and we would like to have more informative labels for the colors and each axis. We can accomplish both of theses goals using scale_color_discrete() and labs().



What if we want the opacity of each point to be proportional to the weight of the vehicle? The alpha parameter lets us control the transparency of a given layer.



What about making the size of the point proportional to the number of cylinders in the vehicle?



Of course, this is getting a bit ridiculous now, but the point is that you can do all sorts of interesting things with ggplot by passing parameters through the layers via the aes() function.

3 Bar Charts

The geom_bar() function allows you to create all sorts of bar charts. We're going to use a different dataset now that comes with ggplot2 called diamonds. This dataset contanis prices and other attributes of more than 50,000 diamonds. The variables included are:

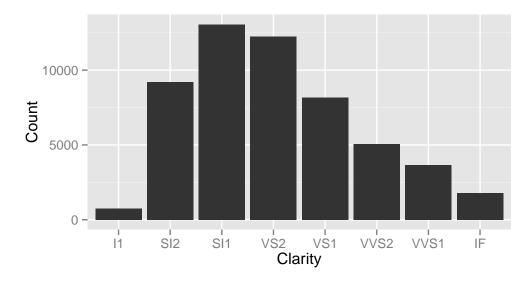
• price: price in US dollars (\$326–\$18,823)

- carat: weight of the diamond (0.2–5.01)
- cut: quality of the cut (Fair, Good, Very Good, Premium, Ideal)
- colour: diamond colour, from J (worst) to D (best)
- clarity: a measurement of how clear the diamond is (I1 (worst), SI1, SI2, VS1, VS2, VVS1, VVS2, IF (best))
- x: length in mm (0–10.74)
- y: width in mm (0–58.9)
- z: depth in mm (0–31.8)
- depth: total depth percentage
- table: width of top of diamond relative to widest point (43–95)

```
head(diamonds)
                 cut color clarity depth table price
##
     carat
## 1 0.23
               Ideal
                         Ε
                               SI2 61.5
                                             55
                                                  326 3.95 3.98 2.43
## 2 0.21
                         Ε
             Premium
                               SI1
                                    59.8
                                             61
                                                  326 3.89 3.84 2.31
## 3
     0.23
                         Ε
                                                  327 4.05 4.07 2.31
                Good
                               VS1
                                   56.9
                                             65
## 4
     0.29
                                                  334 4.20 4.23 2.63
             Premium
                         Ι
                               VS2 62.4
                                             58
## 5
     0.31
                Good
                         J
                               SI2
                                    63.3
                                             58
                                                  335 4.34 4.35 2.75
                         J
## 6 0.24 Very Good
                              VVS2
                                    62.8
                                             57
                                                  336 3.94 3.96 2.48
```

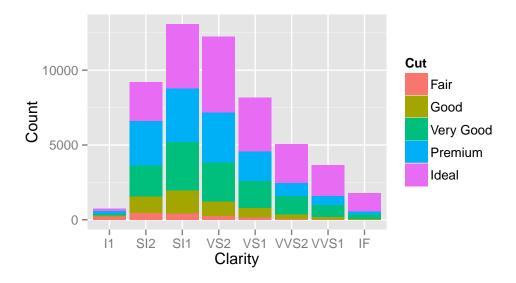
Let's say we want to look at how many diamonds there for each value of clarity. This time we don't need to include a y parameter because the geom_bar() function will use the number of data points in each category as the y value. If you want to pass your own y values then you will need to add stat = "identity" as a parameter in the geom_bar() function.

```
ggplot(data = diamonds, aes(x = clarity)) +
    geom_bar() +
    labs(x = "Clarity", y = "Count")
```



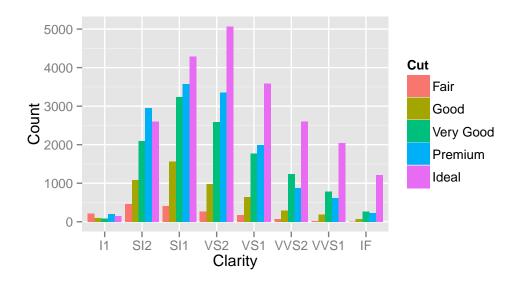
Next, let's color the bars according to the quality of the cut. The fill parameter controls the color of the inside of the bar, and the color parameter controls the color of the outline of the bar. This distinction is used throughout ggplot.

```
ggplot(data = diamonds, aes(x = clarity, fill = cut)) +
    geom_bar() +
    labs(x = "Clarity", y = "Count", fill = "Cut")
```



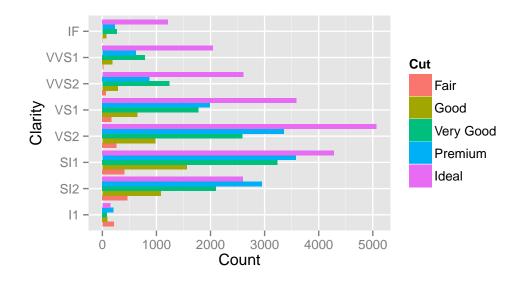
What if we want the bars to be next to each other instead of stacked on top of each other? The position parameter allows us to do that by taking in values of stack, dodge, or fill.

```
ggplot(data = diamonds, aes(x = clarity, fill = cut)) +
    geom_bar(position = "dodge") +
    labs(x = "Clarity", y = "Count", fill = "Cut")
```



Often you will come across text labels for your x-axis that are too large to fit. You can rotate your x-axis labels to make them fit (that's an exercise left to the reader), or you can use coord_flip() to flip the axes.

```
ggplot(data = diamonds, aes(x = clarity, fill = cut)) +
    geom_bar(position = "dodge") +
    labs(x = "Clarity", y = "Count", fill = "Cut") +
    coord_flip()
```



4 Box Plots

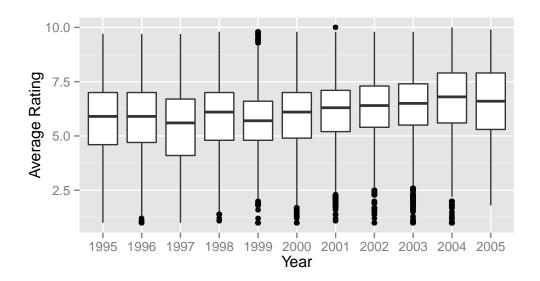
Let's move on to a new dataset called movies. This data comes from IMDB and includes 30000 rows with the following variables:

- title: Title of the movie.
- year: Year of release.
- budget: Total budget (if known) in US dollars
- length: Length in minutes.
- rating: Average IMDB user rating.
- votes: Number of IMDB users who rated this movie.
- r1-10: Multiplying by ten gives percentile (to nearest 10%) of users who rated this movie a 1.
- mpaa: MPAA rating.

• action, animation, comedy, drama, documentary, romance, short: Binary variables representing if movie was classified as belonging to that genre.

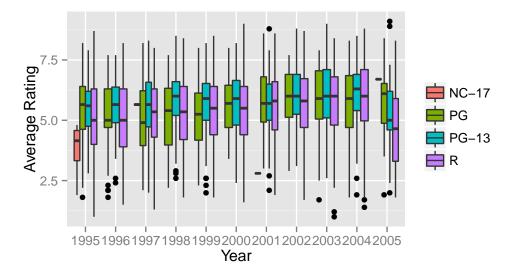
Let's make a boxplot of the average ratings from 1995 to 2005. The geom_boxplot() function allows us to do that.

```
ggplot(data = movies[movies$year >= 1995,] ,
   aes(x = factor(year), y = rating)) +
   geom_boxplot() +
   labs(x = "Year", y = "Average Rating")
```



We can also separate this by the MPAA rating. I'm going to remove the legend title because I feel like it is self explanatory in this context.

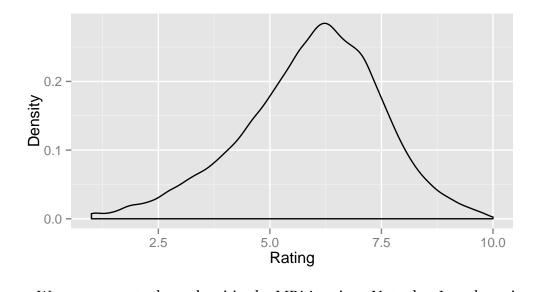
```
ggplot(data = movies[movies$year >= 1995 & movies$mpaa != "",],
    aes(x = factor(year), y = rating, fill = mpaa)) +
    geom_boxplot() +
    labs(x = "Year", y = "Average Rating", fill = "")
```



5 Density

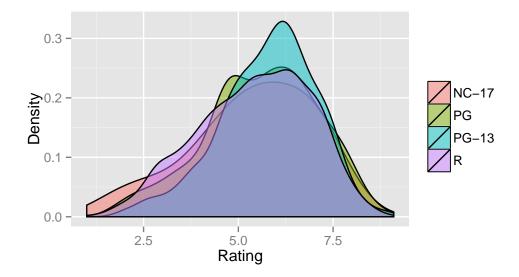
The geom_density() function gives a 1d kernel estimate of whatever variable you give it. The function calls the base R density function which takes a number of parameters. The default is a Gaussian kernel. Let's look at the distribution of movie ratings.

```
ggplot(data = movies, aes(x = rating)) +
    geom_density() +
    labs(x = "Rating", y = "Density")
```



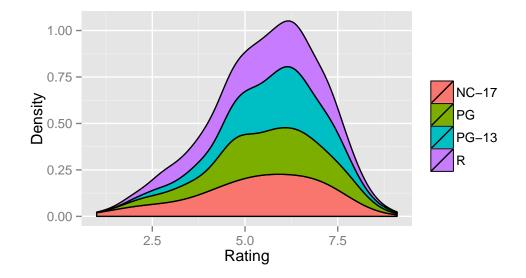
We can separate these densities by MPAA rating. Note that I am lowering the transparency to make it possible to see multiple distributions at once.

```
ggplot(data = movies[movies$mpaa != "",],
  aes(x = rating, fill = mpaa)) +
  geom_density(alpha = 0.5) +
  labs(x = "Rating", y = "Density", fill = "")
```



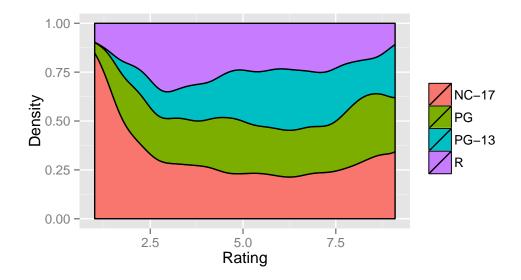
You can also stack these densities on top of each other.

```
ggplot(data = movies[movies$mpaa != "",],
   aes(x = rating, fill = mpaa)) +
   geom_density(position = "stack") +
   labs(x = "Rating", y = "Density", fill = "")
```



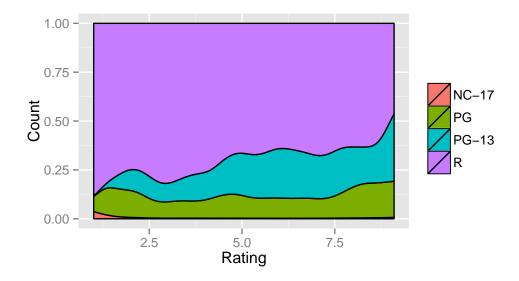
Using position = "fill" can be particularly interesting in this context. That will produce a conditional density estimate for each average rating.

```
ggplot(data = movies[movies$mpaa != "",],
   aes(x = rating, fill = mpaa)) +
   geom_density(position = "fill") +
   labs(x = "Rating", y = "Density", fill = "")
```



Changing it to produce a raw count of records rather than a density estimate might be a better to get an actual idea of how many movies of different MPAA ratings are given a particular average IMDB rating. The special ..count.. expression will do that for us.

```
ggplot(data = movies[movies$mpaa != "",],
   aes(x = rating, y = ..count.., fill = mpaa)) +
   geom_density(position = "fill") +
   labs(x = "Rating", y = "Count", fill = "")
```



6 Smoothing

Sometimes you want to be able to visualize a trend, whether it be a simple linear model or a more complicated smoothing model. The geom_smooth() function takes in several parameters, the most important of which is method which can take the following values:

- lm: linear model
- glm: generalized linear model
- gam: generalized additive model
- loess: LOcal regrESSion (locally weighted scatterplot smoothing)
- rlm: robust linear model

All of these are extensively documented in the R help files. The function defaults to loess for data with <1000 samples and defaults to gam otherwise. For visualization purposes, leaving it to be default will usually work well enough,

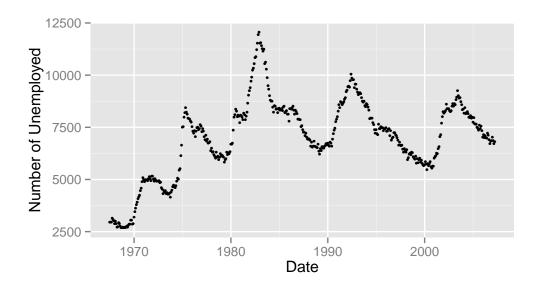
but this function is flexible enough to implement any smoothing method you can think of.

Let's look at yet another new dataset called economics which includes time series data on US economic metrics. The variables included are:

- date. Month of data collection
- psavert, personal savings rate, http://research.stlouisfed.org/fred2/ series/PSAVERT/
- pce, personal consumption expenditures, in billions of dollars, http://research.stlouisfed.org/fred2/series/PCE
- unemploy, number of unemployed in thousands, http://research.stlouisfed. org/fred2/series/UNEMPLOY
- uempmed, median duration of unemployment, in week, http://research.stlouisfed.org/fred2/series/UEMPMED
- pop, total population, in thousands, http://research.stlouisfed.org/ fred2/series/POP

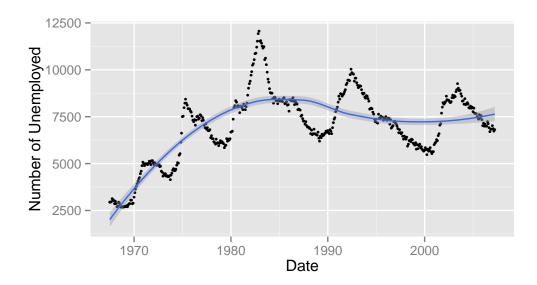
Let's start with something easy like total number of unemployed people over time. First, we have to talk about how dates and times are handled in R. For dates, R has a special variable type called Date. These can be created with the as.Date() function, but luckily for us our dates are already in the correct format. Ggplot has been coded to handle R date and time objects in a way that makes sense, but custom options can be specified using the scale_x_time() function and similar functions.

```
ggplot(data = economics, aes(x = date, y = unemploy)) +
    geom_point(size = 1) +
    labs(x = "Date", y = "Number of Unemployed")
```



Notice how ggplot automatically formats the x-axis in a way that makes sense. Isn't that neat? Next, we'll add a smoothing function to the data to get a better look at what the overall trend is.

```
ggplot(data = economics, aes(x = date, y = unemploy)) +
    geom_point(size = 1) +
    geom_smooth(method = "loess") +
    labs(x = "Date", y = "Number of Unemployed")
```



This is mostly just a proof of concept, in practice you would want to use an actual time series analysis technique for this application (e.g. exponential smoothing, ARIMA).

7 Faceting

What if you want to plot 4 different time series at once, possibly with different scales? The concept of facetting in ggplot allows you to do that. The first step is to transform your data from wide to long format. Wide format is what most datasets look like because it is what makes the most sense to us. Due to the way R handles dataframes, it is often easier to work with long format data from a coding standpoint. Here's what our data currently looks like:

```
head(economics)
##
           date
                          pop psavert uempmed unemploy
                   рсе
## 1 1967-06-30 507.8 198712
                                   9.8
                                           4.5
                                                    2944
   2 1967-07-31 510.9 198911
                                           4.7
                                   9.8
                                                    2945
## 3 1967-08-31 516.7 199113
                                   9.0
                                           4.6
                                                    2958
```

```
## 4 1967-09-30 513.3 199311 9.8 4.9 3143
## 5 1967-10-31 518.5 199498 9.7 4.7 3066
## 6 1967-11-30 526.2 199657 9.4 4.8 3018
```

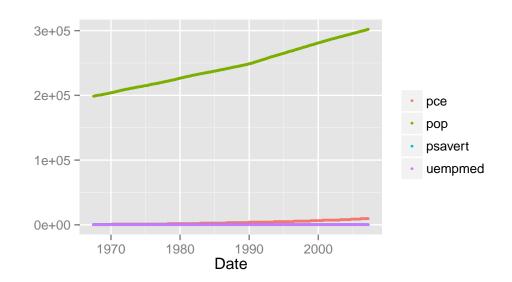
Now we're going to use the melt() function in the reshape2 package to transform this into long form data, using date as an ID variable.

```
library(reshape2)
economicsLong <- melt(economics[,1:5], id = "date")</pre>
head(economicsLong)
           date variable value
##
## 1 1967-06-30
                     pce 507.8
## 2 1967-07-31
                     pce 510.9
## 3 1967-08-31
                     pce 516.7
## 4 1967-09-30
                     pce 513.3
## 5 1967-10-31
                     pce 518.5
## 6 1967-11-30
                     pce 526.2
tail(economicsLong)
##
              date variable value
## 1907 2006-10-31 uempmed
                               8.2
## 1908 2006-11-30 uempmed
                               7.3
## 1909 2006-12-31
                    uempmed
                               8.1
## 1910 2007-01-31
                    uempmed
                               8.1
## 1911 2007-02-28
                    uempmed
                               8.5
## 1912 2007-03-31
                    uempmed
                               8.7
```

Basically, we have collapsed all of our columns (except date) into a single column named value, and we keep track of which row corresponds to which of our original variables in a new column called variable. Let's look at how we

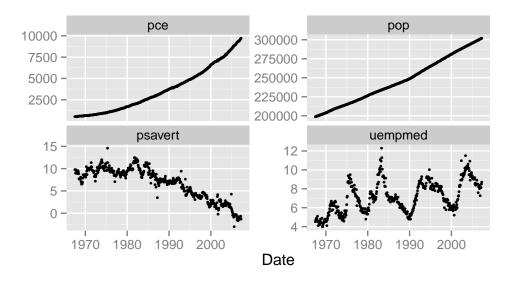
can use this new format to easily create more complicated plots. First, let's plot each variable as a time series on the same plot.

```
ggplot(data = economicsLong, aes(x = date, y = value, color = variable)) +
    geom_point(size = 1) +
    labs(x = "Date", y = "", color = "")
```



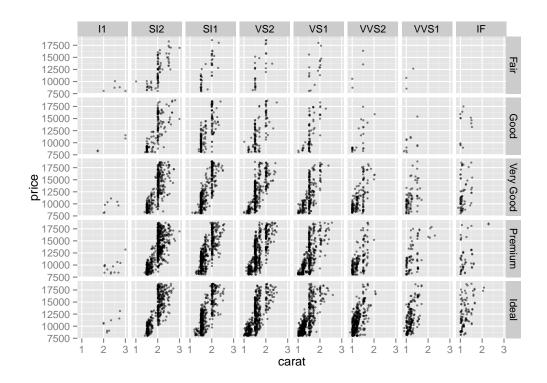
Well, that's obviously not very useful. I want to be able to look at all of my variables at once, but I need them to be on separate plots. This is where the facet_wrap() function comes in handy. Our data is in long format, so it is easy to transform this useless plot into something way more useful.

```
ggplot(data = economicsLong, aes(x = date, y = value)) +
    geom_point(size = 1) +
    facet_wrap(~variable, scale = "free_y") +
    labs(x = "Date", y = "")
```



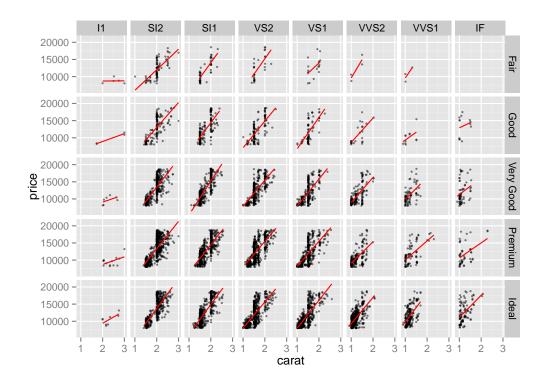
For a better example let's go all the way back to our diamonds data set. Say we want to look at the relationship between carat and price, but now we want to look at how that relationship changes for different cut qualities and clarity rankings. Sounds complicated, but it's actually very easy with the facet_grid() function.

```
library(scales)
ggplot(data = diamonds[diamonds$price > 8000 &
    diamonds$carat <= 3,], aes(x = carat, y = price)) +
    geom_point(size = 1, alpha = 0.5) +
    facet_grid(cut ~ clarity) +
    scale_x_continuous(breaks = pretty_breaks(n = 3))</pre>
```



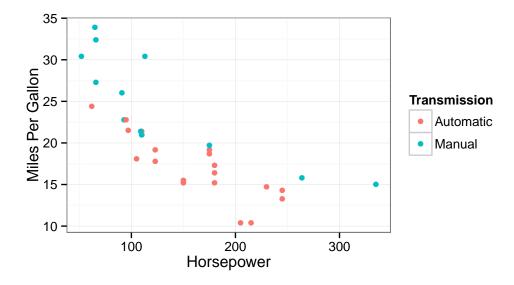
I had to use the pretty_breaks() function in the scales library in order to fix the x-axis because the default choice by ggplot took up too much room. I also subsetted the dataset because plotting 50,000 points in a pdf makes it unnecessarily large. What if we want to add a linear regression over every one of these plots? All we have to do is add geom_smooth().

```
library(scales)
ggplot(data = diamonds[diamonds$price > 8000 &
    diamonds$carat <= 3,], aes(x = carat, y = price)) +
    geom_point(size = 1, alpha = 0.5) +
    geom_smooth(method = "lm", se = FALSE, color = "red") +
    facet_grid(cut ~ clarity) +
    scale_x_continuous(breaks = pretty_breaks(n = 3))</pre>
```

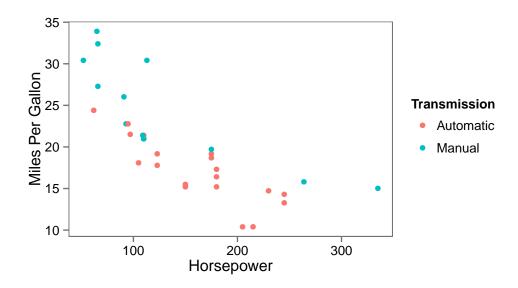


8 Themes and Colors

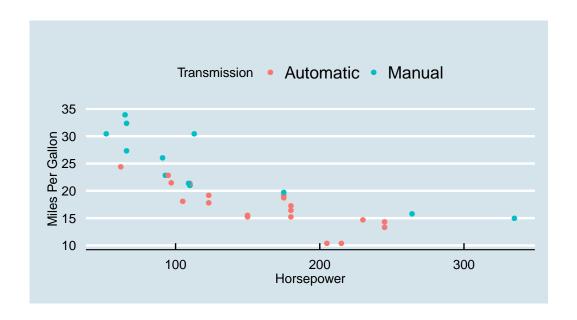
The default theme in ggplot looks pretty nice, but personally I like things to be a bit more minimalist. The theme() function allows you to set all sorts of options for how the overall plot looks, but let's just look at the theme_bw() function.

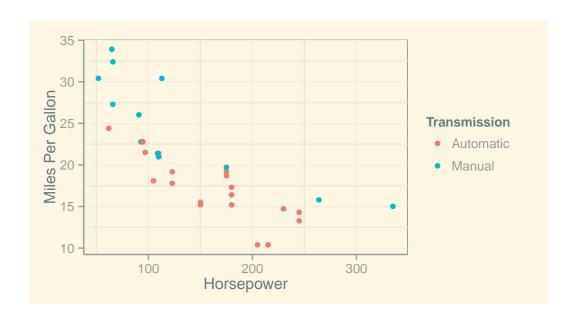


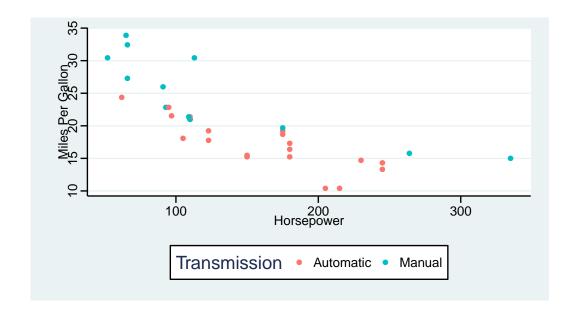
Of course this is mostly personal preference, but I think that looks nicer than the default plot style. The ggplot package comes with many themes already built in, but the ggthemes package offers even more. My personal favorite is theme_few(). If you're a fan of minimalist plots then this is the theme for you.

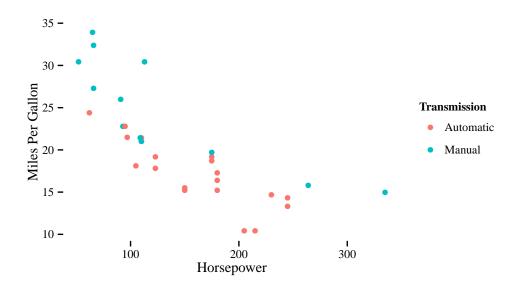


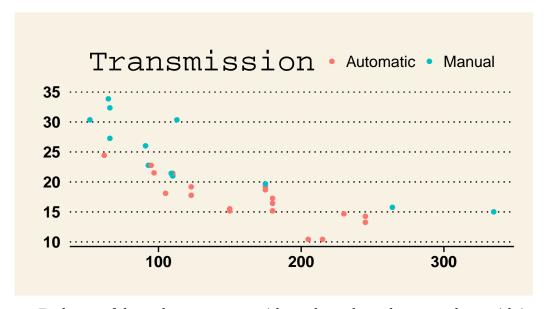
Here are a few more examples:





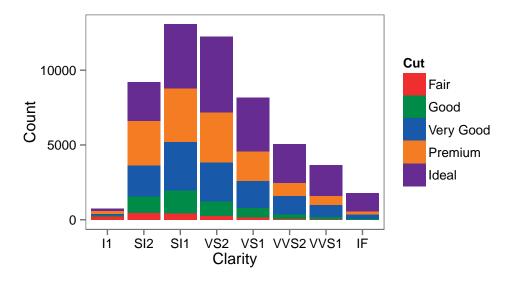






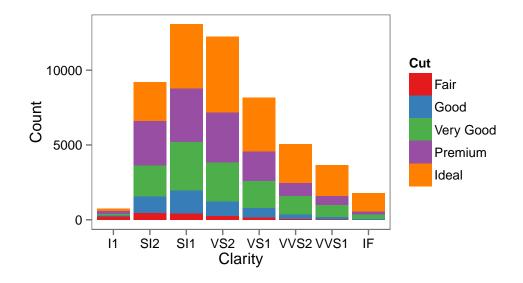
Each one of these themes comes with a color palette that goes along with it. Again, my favorite is the few theme, and you can use the corresponding color palette in the following way:

```
ggplot(data = diamonds, aes(x = clarity, fill = cut)) +
    geom_bar() +
    labs(x = "Clarity", y = "Count", fill = "Cut") +
    theme_few() +
    scale_fill_few(palette = "dark")
```

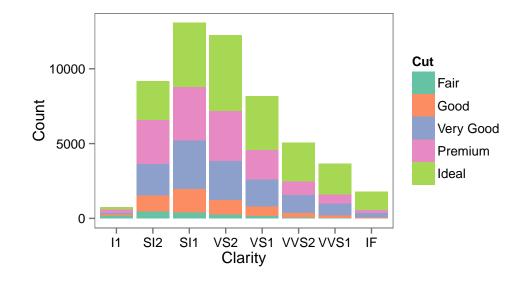


I also feel the need to mention something called Color Brewer (http://colorbrewer2.org/). It's made for discrete values on maps but it can be generalized to other types of plots and even continuous values. It offers a lot of really good color palettes, and it comes built into ggplot (the RColorBrewer package offers access to these palettes outside of ggplot). Here's an example of a few color palettes.

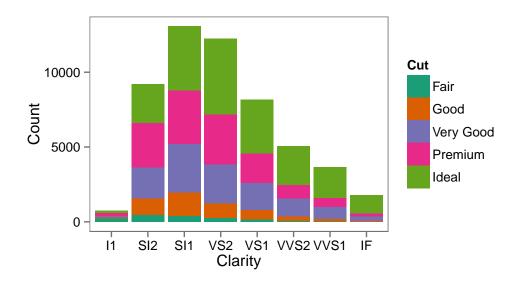
```
ggplot(data = diamonds, aes(x = clarity, fill = cut)) +
    geom_bar() +
    labs(x = "Clarity", y = "Count", fill = "Cut") +
    theme_few() +
    scale_fill_brewer(palette = "Set1")
```



```
ggplot(data = diamonds, aes(x = clarity, fill = cut)) +
    geom_bar() +
    labs(x = "Clarity", y = "Count", fill = "Cut") +
    theme_few() +
    scale_fill_brewer(palette = "Set2")
```



```
ggplot(data = diamonds, aes(x = clarity, fill = cut)) +
    geom_bar() +
    labs(x = "Clarity", y = "Count", fill = "Cut") +
    theme_few() +
    scale_fill_brewer(palette = "Dark2")
```



What about continuous values? Luckily ggplot has provided scale_color_distiller() and similar functions in order to handle this. The following plot colors each point by the average IMDB user rating according to the Reds palette from Color Brewer.

```
ggplot(data = movies[movies$budget >= 1000000,],
    aes(x=budget/1000000, y = length, color = rating)) +
    geom_point(size = 1.5, alpha = 0.5) +
    labs(x = "Budget (million USD)", y = "Length (minutes)",
        color = "Rating") +
    theme_few() +
    scale_color_distiller(palette = "Reds") +
    guides(color = guide_legend(override.aes =
```

list(size = 3, alpha = 1), reverse = TRUE))

